

PCT09

RAW SEQUENCE LISTING

DATE: 01/30/2003

PATENT APPLICATION: US/09/701,500

TIME: 15:50:23

Input Set : A:\TSRI-6511seq.txt

Output Set: N:\CRF4\01302003\I701500.raw

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3 <110> APPLICANT: CHERESH, David A.
         ELICEIRI, Brian
 5
         SCHWARTZBERG, Pamela L.
 7 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS USEFUL FOR MODULATION OF
         ANGIOGENESIS USING TYROSINE KINASE SRC
10 <130> FILE REFERENCE: TSRI 651.1
12 <140> CURRENT APPLICATION NUMBER: US 09/701,500
13 <141> CURRENT FILING DATE: 2000-11-29
15 <150> PRIOR APPLICATION NUMBER: PCT/US99/11780
16 <151> PRIOR FILING DATE: 1999-05-28
18 <150> PRIOR APPLICATION NUMBER: US 60/087,220
19 <151> PRIOR FILING DATE: 1998-05-29
21 <160> NUMBER OF SEQ ID NOS: 6
23 <170> SOFTWARE: PatentIn Ver. 2.0
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38 <223> OTHER INFORMATION: pBR322 sequences
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50 < 223 > OTHER INFORMATION: upstream (numbering begins at the upstream R)
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60 <223> OTHER INFORMATION: R
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63 <221> NAME/KEY: misc_feature

62 <220> FEATURE:



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- 68 <220> FEATURE:
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- 74 <221> NAME/KEY: LTR
- 75 <222> LOCATION: (7166)..(7494)
- 76 <223> OTHER INFORMATION: downstream
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- 79 <221> NAME/KEY: misc_feature
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- 81 <223> OTHER INFORMATION: U3
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- 86 <222> LOCATION: (7394)..(7414)
- 87 <223> OTHER INFORMATION: R
- 90 <220> FEATURE:
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- 93 <223> OTHER INFORMATION: U5
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- 97 <221> NAME/KEY: misc_feature
- 98 <222> LOCATION: (7154)..(7165)
- 99 <223> OTHER INFORMATION: PPT
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- 103 <222> LOCATION: (388)...(391)
- 104 <223> OTHER INFORMATION: splice donor (AGGT)
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- 108 <222> LOCATION: $(507\overline{4})$...(5077)
- 109 <223> OTHER INFORMATION: env splice acceptor (AGGC)
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- 113 <222> LOCATION: (6982)..(6985)
- 114 <223> OTHER INFORMATION: ClaI splice acceptor (AGGA)
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- 125 <222> LOCATION: (909)..(1094)
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- 131 <222> LOCATION: (1095)..(1814)
- 132 <223> OTHER INFORMATION: gag p27



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- 171 <222> LOCATION: (5244)..(6263)
- 172 <223> OTHER INFORMATION: env gp85
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- 178 <222> LOCATION: (6264)..(6878)
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- Dam+ strains and does not cut.
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- 201 atcgggaggc gagttcgatg accetggtgg agggggctgc ggcttaggga ggcagaagct 240
- 203 gagtaccgtc ggagggagct ccagggcccg gagcgactga cccctgccga gaactcagag 300
- ggtcgtcgga agacggagag tgagcccgac gaccacccca ggcacgtctt tggtcggcct 360 205
- gcggatcaag catggaagcc gtcattaagg tgatttcgtc cgcgtgtaaa acctattgcg 420



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213	cgctctccca	ı gcgggcaatg	, gtacttgga <i>a</i>	ı aatcgggaga	gttaaaaacc	tagadattag	600
215	ttttgggggc	: attgaaggcg	, gctcgagagg	, aacaggttac	: atctgagcaa	gcaaagtttt	660
217	ggttgggatt	agggggaggg	, agggtctctc	ccccaggtcc	: ggagtgcatc	gagaaaccag	720
219	clacggagcg	gcgaatcgac	: aaaggggagg	r aggtgggaga	aacaactgtg	cagcgagatg	780
221	cgaagatggc	gccagaggaa	gcggccacac	ctaaaaccqt	tggcacatcc	toctatcatt	840
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242	qqcqqqqqa	acqqactaac	ttggatcgat	taaagggctt	agctgatggg	aacggccaag	1380
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Input Set : A:\TSRI-6511seq.txt

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400	ggggatcgcg	catggcaagg	aattcccagt	cgtccggtag	ggggcccctg	ctatttaggc	6180
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VERIFICATION SUMMARY

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